Exhibit II

Blast 2 Sequences results

PubMed   Entrez   BLAST   OMIM   Taxonomy   Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.13 [Nov-27-2005]

Matrix BLOSUM62 ▾ gap open: 11 gap extension: 1
x_dropoff: 50   expect: 10.000 wordsize: 3   Filter ☑ View option Standard   ▾
Masking character option X for protein, n for nucleotide ▾   Masking color option Black ▾
☐ Show CDS translation   Align

Sequence 1: gi|47117817|sp|O75376|NCOR1_HUMANNuclear receptor corepressor 1 (N-CoR1) (N-CoR).
Length = 296 (736 .. 1031)   SEQ ID NO. 11

Sequence 2: gi|1045655|gb|AAC50236.1|silencing mediator of retinoid and thyroid hormone action
Length = 1495   HSU 37146

No significant similarity was found

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CPU time:      0.02 user secs.        0.02 sys. secs        0.04 total secs.

Lambda      K        H
  0.307     0.125    0.360

Gapped
Lambda      K        H
  0.267     0.0410   0.140


Matrix: BLOSUM62
Gap Penalties: Existence: 11, Extension: 1
Number of Sequences: 1
Number of Hits to DB: 3917
Number of extensions: 2470
Number of successful extensions: 1
Number of sequences better than 10.0: 0
Number of HSP's gapped: 1
Number of HSP's successfully gapped: 0
Length of query: 296
Length of database: 1,196,146,007
Length adjustment: 132
Effective length of query: 164
Effective length of database: 1,196,145,875
Effective search space: 196167923500
Effective search space used: 196167923500
Neighboring words threshold: 9
X1: 16 ( 7.1 bits)
X2: 129 (49.7 bits)
X3: 129 (49.7 bits)
S1: 42 (21.6 bits)
S2: 77 (34.3 bits)
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